GenePattern

SelectFeaturesColumns Documentation

Module name: SelectFeaturesColumns
Description: Takes a column slice from a .res, gct, .odf, or .cls file.
Author: Joshua Gould (Broad Institute), gp-help@broad.mit.edu

Summary: This task generates a new file from a column slice of an existing file. For example, a res file can be produced containing only the tumor samples from a larger res file containing data for both tumor and normal samples. Specific rows can be extracted using the *SelectFeaturesRows* task.

Parameters:

Name	Description
input filename	.res, .gct, .odf, .cls file from which the slice is extracted
columns	the columns to extract (e.g. 0-3, 5, 9)
output	name of the output file

Return Value:
1. a .res, gct, .odf, or .cls file constituted by the specified column slice

Platform dependencies:

Task type:	GeneListSelection
CPU type:	any
OS:	any
Java JVM level:	1.4
Language:	Java